GenCore version 4.5
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OM protein - protein search, using sw model

Run on: May 23, 2001, 14:18:15 ; Search time 18.6 Seconds
(without alignments)
1625.704 Million cell updates/sec

Title: US-08-883-036A-2
Perfect score: 2327
Sequence: 1 MEQRGQNAPAASGARKRHGP.....HLLSSGKFMYLEGNADSAMS 440

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 198801 seqs, 68722935 residues

Total number of hits satisfying chosen parameters: 198801

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : PIR_67:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	243	10.4	461	2	JC4302	tumor necrosis fac
2	218	9.4	454	1	GQMST1	tumor necrosis fac
3	215	9.2	455	1	GQHUT1	tumor necrosis fac
4	209	9.0	435	2	I54182	tumor necrosis fac
5	199.5	8.6	314	2	I37383	FAS soluble protei
6	197	8.5	335	2	A40036	apoptosis-mediatin
7	179.5	7.7	327	2	A46484	apoptosis-mediatin
8	173.5	7.5	461	1	GQRTT1	tumor necrosis fac
9	172	7.4	461	1	A35356	tumor necrosis fac
10	167.5	7.2	459	2	I48854	gene murine tumour
11	166	7.1	427	1	GQHUN	nerve growth facto
12	163.5	7.0	474	2	B38634	tumor necrosis fac
13	161.5	6.9	416	1	JN0006	nerve growth facto
14	155	6.7	425	1	A26431	nerve growth facto
15	148.5	6.4	324	2	JC2395	Fas antigen precur
16	144	6.2	348	2	T28623	hypothetical prote
17	142	6.1	272	2	I48700	gene ox40 protein
18	141	6.1	349	2	D36858	gene G4R protein -
19	140	6.0	349	2	D72175	G2R protein - vari
20	139.5	6.0	595	2	A42086	CD30 antigen precu
21	137	5.9	271	2	S12783	OX40 antigen precu
22	136.5	5.9	277	2	A60771	B-cell activation
23	132	5.7	557	2	A48434	variant-specific s
24	131.5	5.7	577	2	A60501	thrombomodulin pre
25	129.5	5.6	277	2	I37552	OX40 homolog - hum
26	128	5.5	326	1	GQVZML	T2 protein - myxom
27	128	5.5	2824	2	T22759	hypothetical prote
28	126	5.4	256	2	B32393	T-cell antigen 4-1
29	124.5	5.4	521	2	A29345	steroid hormone re
30	124.5	5.4	932	2	A42632	cell adhesion mole
31	123	5.3	493	2	JC5486	membrane glycoprot
32	122.5	5.3	1372	2	T25933	hypothetical prote
33	119	5.1	1036	2	T17405	scavenger receptor
34	118.5	5.1	305	2	A46476	B cell-associated
35	118.5	5.1	356	2	A25918	thrombomodulin - b
36	116.5	5.0	575	1	THHUB	thrombomodulin pre
37	116	5.0	984	1	A34076	protein-tyrosine k
38	113	4.9	325	2	B43692	T2 protein - rabbi
39	113	4.9	1810	1	A32230	tenascin precursor
40	112	4.8	1436	2	A46496	antigen WC1.1 prec
41	111.5	4.8	667	2	A48579	trophozoite surfac
42	110.5	4.7	655	1	A46688	hepatocyte growth
43	110.5	4.7	1548	2	S34583	serine proteinase
44	110	4.7	250	1	A49053	CD27 antigen precu
45	109.5	4.7	1766	2	A42125	trophozoite cystei

ALIGNMENTS

RESULT 1
JC4302
tumor necrosis factor receptor p55 precursor - pig
C;Species: Sus scrofa domestica (domestic pig)
C;Date: 29-Nov-1995 #sequence_revision 08-Feb-1996 #text_change 23-Jul-1999
C;Accession: JC4302; PC4093
R;Suter, B.; Pauli, U.
Gene 163, 263-266, 1995
A;Title: Cloning of the cDNA encoding the porcine p55 tumor necrosis factor receptor.
A;Reference number: JC4302; MUID:96011645
A;Accession: JC4302
A;Molecule type: mRNA
A;Residues: 1-461 <SUT>
A;Cross-references: GB:U19994; NID:g1141752; PIDN:AAC48499.1; PID:g1141753
A;Accession: PC4093
A;Molecule type: protein
A;Residues: 1-7 <SU2>
A;Experimental source: kidney cell line 15
C;Genetics:
A;Gene: tnfr
C;Superfamily: tumor necrosis factor receptor type 1; NGF receptor repeat homology
C;Keywords: glycoprotein; kidney; receptor; transmembrane protein; tumor
F;1-29/Domain: signal sequence #status predicted <SIG>
F;30-461/Product: tumor necrosis factor receptor p55 #status predicted <MAT>
F;44-194/Domain: extracellular cysteine rich #status predicted <EXT>
F;44-82/Domain: NGF receptor repeat homology <NG1>
F;84-126/Domain: NGF receptor repeat homology <NGF>
F;211-231/Domain: transmembrane #status predicted <TMM>
F;361-447/Domain: signal transduction #status predicted <SIT>
F;54,145,151/Binding site: carbohydrate (Asn) (covalent) #status predicted

Query Match 10.4%; Score 243; DB 2; Length 461;
Best Local Similarity 24.6%; Pred. No. 7.5e-10;
Matches 119; Conservative 61; Mismatches 185; Indels 118; Gaps 26;

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Qy      30 PGPRVPKTLVLVVAAVLLLV---SAESALITQQDLAPQQRAAPQQKRSSPSE-----GLC 81
           ||  :|    ||: |:|: |        ::   |   ::   || | | |         |
Db       7 PGLLLP----LVLRALLVDVYPAGVHGLVLHPGDREKRESLCPQGKYSHPQNRSICCTKC 62

Qy      82 PPGHHISED------GRDCISCKYGQDYSTHWNDLLFCLRCTRCDS--GEVELSPCTTTR 133
             | ::  |        ||  |  |  ::   | |  || |::| |   :||:|||  |
Db      63 HKGTYLHNDCLGPGLDTDCRECDNG-TFTASENHLTQCLSCSKCRSEMSQVEISPCTVDR 121

Qy     134 NTVCQCEEGTFREEDSPEM--CRKCRTGCPRGMVKVGDCTPWSD---------------- 175
           :||| | :  :|:  |  :  |  |   || | |::  |   |
Db     122 DTVCGCRKNQYRKYWSETLFQCLNCSL-CPNGTVQL-PCLEKQDTICNCHSGFFLRDKEC 179

Qy     176 IECVH-KESGTKHSGEAPAVEETVTSSPGTPASPCSLSGIIIGVTVAAVVLIVAVFVCKS 234
           : ||: | :  |:    ||  ||       |  :       |  |: :|  : :     |:
Db     180 VSCVNCKNADCKNL--CPATSETRNDFQDTGTTVLLPLVIFFGLCLAFFLFV--GLACRY 235
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Qy    235 LLWKKVLPYLKGICSGGG-----GDPERVDRS-------------SQRPGAEDNVLNEIV 276
Db    236 QRWK---PKLYSIICGKSTPVKEGEPEPLATAPSFGPITTFSPIPSFSPTTTFSPVPSFS 292

Qy    277 SILQPTQVP--EQEMEVQEPAE-----PTGVNMLSP---------------GESEHLLEP 314
Db    293 PISSPTFTPCDWSNIKVTSPPKEIAPPPQGAGPILPMPPASTPVPTPLPKWGGSAHSAHS 352

Qy    315 AEAERSQRRRLLVPANEGDPTETLRQCFDDFADLVPFDSWEPLMRKLGLMDNEI-KVAKA 373
Db    353 APAQLA----------DADPA-TLYAVVDG----VPPTRWKEFVRRLGLSEHEIERLELQ 397

Qy    374 EAAGHRDTLYTMLIKWVNKTG-RDASVH---------TLLDALETLGERL---AKQKIED 420
Db    398 NGRCLREAQYSMLAEWRRRTSRREATLELLGSVLRDMDLLGCLEDIEEALRGPARLAPAP 457

Qy    421 HLL 423
Db    458 HLL 460
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RESULT 2
GQMST1
tumor necrosis factor receptor 1 precursor - mouse
N;Alternate names: tumor necrosis factor receptor, 55K
C;Species: Mus musculus (house mouse)
C;Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 01-Dec-2000
C;Accession: A38634; B40254; S16677; S19021; I54532; I57826
R;Lewis, M.; Tartaglia, L.A.; Lee, A.; Bennett, G.L.; Rice, G.C.; Wong, G.H.W.; Chen, E.
Proc. Natl. Acad. Sci. U.S.A. 88, 2830-2834, 1991
A;Title: Cloning and expression of cDNAs for two distinct murine tumor necrosis factor r
A;Reference number: A38634; MUID:91187885
A;Accession: A38634
A;Molecule type: mRNA
A;Residues: 1-454 <LEW>
A;Cross-references: GB:M60468; NID:g199825; PIDN:AAA39751.1; PID:g199826
R;Goodwin, R.G.; Anderson, D.; Jerzy, R.; Davis, T.; Brannan, C.I.; Copeland, N.G.; Jenk
Mol. Cell. Biol. 11, 3020-3026, 1991
A;Title: Molecular cloning and expression of the type 1 and type 2 murine receptors for
A;Reference number: A40254; MUID:91246168
A;Accession: B40254
A;Molecule type: mRNA
A;Residues: 1-454 <GO2>
A;Cross-references: GB:M60468; NID:g199825; PIDN:AAA39751.1; PID:g199826
R;Barrett, K.; Taylor-Fishwick, D.A.; Cope, A.P.; Kissonerghis, A.M.; Gray, P.W.; Feldma
Eur. J. Immunol. 21, 1649-1656, 1991
A;Title: Cloning, expression and cross-linking analysis of the murine p55 tumor necrosis
A;Reference number: S16677; MUID:91285014
A;Accession: S16677
A;Molecule type: mRNA
A;Residues: 1-454 <BAR>
A;Cross-references: EMBL:X59238; NID:g53578; PIDN:CAA41922.1; PID:g53579
R;Rothe, J.G.; Brockhaus, M.; Gentz, R.; Lesslauer, W.
Immunogenetics 34, 338-340, 1991
A;Title: Molecular cloning and expression of the mouse Tnf receptor type b.
A;Reference number: S19021; MUID:92039815
A;Accession: S19021
A;Molecule type: mRNA
A;Residues: 1-454 <ROT>
A;Cross-references: EMBL:X57796; NID:g54848; PIDN:CAA40936.1; PID:g54849
R;Bebo, B.F.
Immunogenetics 39, 450-451, 1994
A;Title: Nucleotide sequence of the TNF type I receptor from a mouse endothelioma cell l
A;Reference number: I54532; MUID:94245292
A;Accession: I54532
A;Status: translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-454 <RES>
A;Cross-references: GB:L26349; NID:g430732; PIDN:AAA59361.1; PID:g430733
R;Rothe, J.G.; Bluethmann, H.; Gentz, R.; Lesslauer, W.; Steinmetz, M.
Mol. Immunol. 30, 165-176, 1993
A;Title: Genomic organization and promoter function of the murine tumor necrosis factor
A;Reference number: I57826; MUID:93156721
A;Accession: I57826
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-393,'G',395-454 <RE2>
A;Cross-references: GB:M76656; NID:g202100; PIDN:AAA40465.1; PID:g202102
C;Comment: This protein is one of two distantly related receptors for both TNF-alpha
C;Genetics:
A;Gene: TNFR-2
A;Introns: 13/3; 65/1; 108/1; 158/1; 184/2; 210/1; 248/1; 257/3; 353/1
C;Superfamily: tumor necrosis factor receptor type 1; NGF receptor repeat homology
C;Keywords: cytokine receptor; duplication; glycoprotein; receptor; transmembrane pro
F;1-29/Domain: signal sequence #status predicted <SIG>
F;30-454/Product: tumor necrosis factor receptor type 1 #status predicted <MAT>
F;30-212/Domain: extracellular #status predicted <EXT>
F;44-82/Domain: NGF receptor repeat homology <NG1>
F;84-126/Domain: NGF receptor repeat homology <NG2>
F;127-167/Domain: NGF receptor repeat homology <NG3>
F;168-204/Domain: NGF receptor repeat homology <NG4>
F;213-235/Domain: transmembrane #status predicted <MEM>
F;236-454/Domain: intracellular #status predicted <INT>
F;54,151,202/Binding site: carbohydrate (Asn) (covalent) #status predicted

Query Match 9.4%; Score 218; DB 1; Length 454;
Best Local Similarity 22.2%; Pred. No. 4.4e-08;
Matches 109; Conservative 51; Mismatches 163; Indels 168; Gaps 22;

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Qy     32 PRVPKTLVLVVAAVLLLVSAESALITQQDLAPQQRAAPQQKRSSPSEGLCPPGHHI---- 87
Db      4 PTVPGLLLSLVLLALLMGIHPSGV---TGLVPS--LGDREKRDS----LCPQGKYVHSKN 54

Qy     88 ------------------SEDGRD--CISCKYGQDYSTHWNDLLFCLRCTRC--DSGEVE 125
Db     55 NSICCTKCHKGTYLVSDCPSPGRDTVCRECEKG-TFTASQNYLRQCLSCKTCRKEMSQVE 113

Qy    126 LSPCTTTRNTVCQCEEGTFREEDSPEM-----CRKCRTG--------------CPRG-M 164
Db    114 ISPCQADKDTVCGCKENQFQRYLSETHFQCVDCSPCFNGTVTIPCKETQNTVCNCHAGFF 173

Qy    165 VKVGDCTPWSDIECVHKESGTKHSGEAPAVEETVTSSPGTPASPCSLSGIIIGVTVAAVV 224
Db    174 LRESECVPCS--HCKKNEECMKLCLPPPLANVTNPQDSGTAV----LLPLVILLGLCLLS 227

Qy    225 LIVAVFVCKSLLWKKVLPYLKGICSGGGGDPERVDRSSQRPGAEDNVLNEIVSILQPTQV 284
Db    228 FIFISLMCRYPRW-------------------------RP--------EVYSIICRDPV 253

Qy    285 PEQEMEVQEPAEP---------TGVN----MLSPGESE---------------HLLEPA 315
Db    254 PVKEEKAGKPLTPAPSPAFSPTSGFNPTLGFSTPGFSSPVSSTPISPIFGPSNWHFMPPV 313

Qy    316 EAERSQRRRLLVPANEGD----------PTETLRQCFDDFA----------------DL 348
Db    314 SE--------VVPTQGADPLLYESLCSVPAPTSVQKWEDSAHPQRPDNADLAILYAVVDG 365

Qy    349 VPFDSWEPLMRKLGLMDNEI-KVAKAEAAGHRDTLYTMLIKWVNKTGRDASVHTLLDALE 407
Db    366 VPPARWKEFMRFMGLSEHEIERLEMQNGRCLREAQYSMLEAWRRRTPRHE------DTLE 419

Qy    408 TLGERLAKQKI 418
Db    420 VVGLVLSKMNL 430
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RESULT 3
GQHUT1
tumor necrosis factor receptor 1 precursor [validated] - human
N;Alternate names: P55 tumor necrosis factor receptor; TNF receptor type 1
N;Contains: tumor necrosis factor alpha inhibitor; tumor necrosis factor binding prot
C;Species: Homo sapiens (man)
C;Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 08-Dec-2000
C;Accession: A38208; A34899; A34900; A36555; C36555; A38281; S12057; JT0758; A60231;
R;Fuchs, P.; Strehl, S.; Dworzak, M.; Himmler, A.; Ambros, P.F.

Genomics 13, 219-224, 1992
A;Title: Structure of the human TNF receptor 1 (p60) gene (TNRF1) and localization to ch
A;Reference number: A38208; MUID:92250049
A;Accession: A38208
A;Molecule type: DNA
A;Residues: 1-455 <FUC>
A;Cross-references: GB:M75864; GB:M75865; GB:M75866; NID:g339748; PIDN:AAA61201.1; PID:g
R;Loetscher, H.; Pan, Y.C.E.; Lahm, H.W.; Gentz, R.; Brockhaus, M.; Tabuchi, H.; Lesslau
Cell 61, 351-359, 1990
A;Title: Molecular cloning and expression of the human 55 kd tumor necrosis factor recep
A;Reference number: A34899; MUID:90235284
A;Accession: A34899
A;Molecule type: mRNA
A;Residues: 1-455 <LOE>
A;Cross-references: GB:M58286; GB:M33480; NID:g339753; PIDN:AAA36753.1; PID:g339754
A;Experimental source: placenta
A;Note: part of this sequence, including the amino end of the mature protein, confirmed
R;Schall, T.J.; Lewis, M.; Koller, K.J.; Lee, A.; Rice, G.C.; Wong, G.H.W.; Gatanaga, T.
Cell 61, 361-370, 1990
A;Title: Molecular cloning and expression of a receptor for human tumor necrosis factor.
A;Reference number: A34900; MUID:90235285
A;Accession: A34900
A;Molecule type: mRNA
A;Residues: 1-455 <SCH>
A;Cross-references: GB:M33294; NID:g339744; PIDN:AAA03210.1; PID:g339745
R;Himmler, A.; Maurer-Fogy, I.; Kroenke, M.; Scheurich, P.; Pfizenmaier, K.; Lantz, M.;
DNA Cell Biol. 9, 705-715, 1990
A;Title: Molecular cloning and expression of human and rat tumor necrosis factor recepto
A;Reference number: A36555; MUID:91090841
A;Accession: A36555
A;Molecule type: mRNA
A;Residues: 1-455 <HIM>
A;Cross-references: GB:M63121; NID:g339755; PIDN:AAA36754.1; PID:g339756
A;Accession: C36555
A;Molecule type: protein
A;Residues: 30-38;41-53,'X',55-79,'XX',82-94,'NK';'XX',100-104;107-128;162-167,'X',169-2
A;Note: the purified protein, called tumor necrosis factor binding protein, is a soluble
R;Gray, P.W.; Barrett, K.; Chantry, D.; Turner, M.; Feldmann, M.
Proc. Natl. Acad. Sci. U.S.A. 87, 7380-7384, 1990
A;Title: Cloning of human tumor necrosis factor (TNF) receptor cDNA and expression of re
A;Reference number: A38281; MUID:91017509
A;Accession: A38281
A;Molecule type: mRNA
A;Residues: 1-455 <GRA>
A;Cross-references: GB:M37764
A;Note: the authors translated the codon TGG for residue 371 as Thr, AAG for residue 372
R;Nophar, Y.; Kemper, O.; Brakebusch, C.; Engelmann, H.; Zwang, R.; Aderka, D.; Holtmann
EMBO J. 9, 3269-3278, 1990
A;Title: Soluble forms of tumor necrosis factor receptors (TNF-Rs). The cDNA for the typ
le form of the receptor.
A;Reference number: S12057; MUID:91006021
A;Accession: S12057
A;Molecule type: mRNA
A;Residues: 1-455 <NOP>
A;Cross-references: EMBL:X55313; NID:g37223; PIDN:CAA39021.1; PID:g37224
A;Note: parts of soluble TNF binding protein 1, including its amino and carboxyl ends, w
R;Kemper, O.; Wallach, D.
Gene 134, 209-216, 1993
A;Title: Cloning and partial characterization of the promoter for the human p55 tumor ne
A;Reference number: JT0758; MUID:94085779
A;Accession: JT0758
A;Molecule type: DNA
A;Residues: 1-13 <KEM>
R;Seckinger, P.; Vey, E.; Turcatti, G.; Wingfield, P.; Dayer, J.M.
Eur. J. Immunol. 20, 1167-1174, 1990
A;Title: Tumor necrosis factor inhibitor: purification, NH-2-terminal amino acid sequenc
A;Reference number: A60231; MUID:90292116
A;Accession: A60231
A;Molecule type: protein
A;Residues: 41-43,'X',45-53,'X',55-57 <SEC>
R;Gatanaga, T.; Hwang, C.; Kohr, W.; Cappuccini, F.; Lucci III, J.A.; Jeffes, E.W.B.; Le
Proc. Natl. Acad. Sci. U.S.A. 87, 8781-8784, 1990
A;Title: Purification and characterization of an inhibitor (soluble tumor necrosis factd
tients.
A;Reference number: A38258; MUID:91062364
A;Accession: A38258
A;Molecule type: protein
A;Residues: 41-60 <GAT>
A;Experimental source: cancer patient serum
R;Olsson, I.; Lantz, M.; Nilsson, E.; Peetre, C.; Thysell, H.; Grubb, A.; Adolf, G.
Eur. J. Haematol. 42, 270-275, 1989
A;Title: Isolation and characterization of a tumor necrosis factor binding protein fr
A;Reference number: A60594; MUID:89171156
A;Accession: A60594
A;Molecule type: protein
A;Residues: 41-43,'X',45-53,'V',55-57,'XK',60 <OLS>
A;Experimental source: renal failure patient urine
R;Engelmann, H.; Novick, D.; Wallach, D.
J. Biol. Chem. 265, 1531-1536, 1990
A;Title: Two tumor necrosis factor-binding proteins purified from human urine. Eviden
A;Reference number: A35010; MUID:90110215
A;Accession: A35010
A;Molecule type: protein
A;Residues: 41-45 <ENG>
A;Experimental source: normal urine
R;Kajihara, J.; Asada, A.; Kirihara, S.; Kato, K.
Biosci. Biotechnol. Biochem. 58, 2266-2268, 1994
A;Title: Amino acid sequence of natural tumor necrosis factor alpha inhibitor purifie
A;Reference number: JC2404; MUID:95128033
A;Accession: JC2404
A;Molecule type: protein
A;Residues: 41-53,'X',55-144,'X',146-150,'X',152-186,'X',188-201 <KAJ>
A;Experimental source: urine
C;Comment: This protein is one of two known receptors for both TNF-alpha (cachectin)
C;Genetics:
A;Gene: GDB:TNFR1
A;Cross-references: GDB:125913; OMIM:191190
A;Map position: 12p13.2-12p13.2
A;Introns: 13/3; 65/1; 108/1; 158/1; 184/2; 209/1; 247/1; 256/3; 353/1
C;Superfamily: tumor necrosis factor receptor type 1; NGF receptor repeat homology
C;Keywords: duplication; glycoprotein; receptor; transmembrane protein
F;1-21/Domain: signal sequence #status predicted <SIG>
F;22-455/Product: tumor necrosis factor receptor 1 #status predicted <MAT>
F;30-211/Domain: extracellular #status predicted <EXT>
F;41-201/Product: TNF binding protein 1 (tumor necrosis factor alpha inhibitor) #stat
F;44-82/Domain: NGF receptor repeat homology <NG1>
F;84-126/Domain: NGF receptor repeat homology <NG2>
F;127-167/Domain: NGF receptor repeat homology <NG3>
F;168-196/Domain: NGF receptor repeat homology <NG4>
F;212-234/Domain: transmembrane #status predicted <MEM>
F;235-455/Domain: intracellular #status predicted <INT>
F;54,145,151/Binding site: carbohydrate (Asn) (covalent) #status predicted

Query Match 9.2%; Score 215; DB 1; Length 455;
Best Local Similarity 23.0%; Pred. No. 7.2e-08;
Matches 109; Conservative 62; Mismatches 172; Indels 130; Gaps 22;

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Qy      34 VPKTLVLVVAAVLLLVSAESALITQQDLAPQQRAAPQQKRSSPSEGLCPPGHHISEDGR- 92
           ||  |: :|   ||:    | :|    | |       ::|| |    :|| | :|
Db       6 VPDLLLPLVLLELLVGIYPSGVI---GLVP--HLGDREKRDS----VCPQGKYIHPQNNS 56

Qy      93 ----------------------DCISCKYGQDYSTHWNDLLFCLRCTRC--DSGEVELS 127
                                  ||  |: |  ::   | |  || |::|  : |:||:|
Db      57 ICCTKCHKGTYLYNDCPGPGQDTDCRECESG-SFTASENHLRHCLSCSKCRKEMGQVEIS 115

Qy     128 PCTTTRNTVCQCEEGTFREEDSPEM--CRKCRTGCPRGMVKVGDCTPWSDIECVHKESG- 184
           ||  |:||| | :  :|   | :  | |  | | |          : |  |::
Db     116 SCTVDRDTVCGCRKNQYRHYWSENLFQCFNCSL-CLNGTV---------HLSCQEKQNTV 165

Qy     185 -TKHSGEAPAVEETVTSSPGTPASPC---------SLSGIIIGVTVAAVVLIVAVFVC-K 233
            | |:|      | |: |    :  |          :: |     |   : |::   :|
Db     166 CTCHAGFFLRENECVSCSNCKKSLECTKLCLPQIENVKGTEDSGTTVLLPLVIFFGLCLL 225

Qy     234 SLL----------WKKVLPYLKGICSGGGGDPER---VDRSSQRPGAEDNVLNEIVSILQ 280
           |||          ||  |  :  :|  |   ||:   :: :: :| | :   :
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```
Db    226 SLLFIGLMYRYQRWKSKLYSI--VC--GKSTPEKEGELEGTTTKPLAPNPSFSP-----T 276

Qy    281 PTQVPEQEMEVQEPAEPTGVNMLSPGESEHLLEPAEAERSQRRRLLVPANEG-------- 332
Db    277 PGFTPTLGFSPVPSSTFTSSSTYTPGDCPNFAAP--------RREVAPPYQGADPILATA 328

Qy    333 ---DPTETLRQCFDDFA-----------------DLVPFDSWEPLMRKLGLMDNEI-KV 370
Db    329 LASDPIPNPLQKWEDSAHKPQSLDTDDPATLYAVVENVPPLRWKEFVRRLGLSDHEIDRL 388

Qy    371 AKAEAAGHRDTLYTMLIKWVNKT---------GRDASVHTLLDALETLGERL 413
Db    389 ELQNGRCLREAQYSMLATWRRRTPRREATLELLGRVLRDMDLLGCLEDIEEAL 441
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RESULT 4
I54182
tumor necrosis factor receptor 2-related protein - human
C;Species: Homo sapiens (man)
C;Date: 24-May-1996 #sequence_revision 24-May-1996 #text_change 17-Mar-2000
C;Accession: I54182
R;Baens, M.; Chaffanet, M.; Cassiman, J.J.; Van den Berghe, H.; Marynen, P.
Genomics 16, 214-218, 1993
A;Title: Construction and evaluation of a hncDNA library of human 12p transcribed sequen
A;Reference number: I54182; MUID:93252381
A;Accession: I54182
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-435 <RES>
A;Cross-references: GB:L04270; NID:g339761; PIDN:AAA36757.1; PID:g339762
C;Genetics:
A;Gene: GDB:LTBR
A;Cross-references: GDB:1230195; OMIM:600979
A;Map position: 12p13.3-12p13.1
C;Superfamily: tumor necrosis factor receptor type 1; NGF receptor repeat homology

Query Match 9.0%; Score 209; DB 2; Length 435;
Best Local Similarity 24.5%; Pred. No. 1.8e-07;
Matches 91; Conservative 39; Mismatches 129; Indels 112; Gaps 18;

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Qy     20 PGPREARGARPGPRVPKTLVLVVAAVLLLVSAESALITQQDLAPQQRAAPQQKRSSPSE- 78
Db      4 PWATSAPGLAWGP-------LVLGLFGLLAA------SQPQAVPPYASENQTCRDQEKEY 50

Qy     79 ---------GLCPPGHHISEDG---RD--CISCKYGQDYSTHWNDLLFCLRCTRCD--SG 122
Db     51 YEPQHRICCSRCPPGTYVSAKCSRIRDTVCATCA-ENSYNEHWNYLTICQLCRPCDPVMG 109

Qy    123 EVELSPCTTTRNTVCQCEEGTFREEDSPEMCRKCRTGCPRGMVKVGDCTPWSDIE----- 177
Db    110 LEEIAPCTSKRKTQCRCQPGMFCAAWALE-CTHCEL--------LSDCPPGTEAELKDEV 160

Qy    178 ------CVHKESGTKHSGEAPAV----------EETVTSSPGTPAS----------PCS 210
Db    161 GKGNNHCVPCKAGHFQNTSSPSARCQPHTRCENQGLVEAAPGTAQSDTTCKNPLEPLPPE 220

Qy    211 LSGIIIGVTV----AAVVLIVAVFVCKSLLWK-------KVLPYLKGICSGGGGDPERVD 259
Db    221 MSGTMLMLAVLLPLAFFLLLATVFSC---IWKSHPSLCRKLGSLLKRRPQGEGPNPV--- 274

Qy    260 RSSQRPGAEDNVLNEIVSILQPTQ--------------------VPEQEMEVQEPAEPTGV 300
Db    275 AGSWEPPKAHPYFPDLVQPLLPISGDVSPVSTGLPAAPVLEAGVPQQ----QSPLDLTRE 330

Qy    301 NMLSPGESEHL 311
Db    331 PQLEPGEQSQV 341
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RESULT 5
I37383
FAS soluble protein - human
C;Species: Homo sapiens (man)
C;Date: 02-Jul-1996 #sequence_revision 02-Jul-1996 #text_change 21-Jul-2000
C;Accession: I37383
R;Cascino, I.; Fiucci, G.; Papoff, G.; Ruberti, G.
J. Immunol. 154, 2706-2713, 1995
A;Title: Three functional soluble forms of the human apoptosis-inducing Fas molecule
A;Reference number: I37383; MUID:95181785
A;Accession: I37383
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-314 <RES>
A;Cross-references: EMBL:Z47993; NID:g728578; PIDN:CAA88031.1; PID:g695539

Query Match 8.6%; Score 199.5; DB 2; Length 314;
Best Local Similarity 27.7%; Pred. No. 5.9e-07;
Matches 74; Conservative 36; Mismatches 96; Indels 61; Gaps 13;

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Qy     38 LVLVVAAVLLL----VSAESALITQQDLAPQQRAAPQQKRSSPSEGL----------CPP 83
Db      8 LPLVLTSVARLSSKSVNAQVTDINSKGL--ELRKTVTTVETQNLEGLHHDGQFCHKPCPP 65

Qy     84 GHH------ISEDGRDCISCKYGQDYSTHWNDLLFCLRCTRCDSG---EVELSPCTTTRN 134
Db     66 GERKARDCTVNGDEPDCVPCQEGKEYTDKAHFSSKCRRCRLCDEGHGLEVEIN-CTRTQN 124

Qy    135 TVCQCEEGTFREEDSPEMCRKCRTGCPRGMVKVGDCTPWSDIECVH-------KESGTKH 187
Db    125 TKCRCKPNFFCNSTVCEHCDPC-TKCEHGIIK--ECTLTSNTKCKEEVKRKEVQKTCRKH 181

Qy    188 SGEAPAVEETVTSSPGTPA---SPCSLSGIIIGVTVAAVVLIVAVFVCKSLLWKKVLPYL 244
Db    182 RKENQGSHESPTLNPETVAINLSDVDLSKYI--TTIAGVM--------------TLSQV 224

Qy    245 KGICSGGGGDPERVDRSSQRPGAEDNV 271
Db    225 KGFVRKNGVNEAKIDEIK-----NDNV 246
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RESULT 6
A40036
apoptosis-mediating surface antigen Fas precursor - human
N;Alternate names: surface antigen APO-1
C;Species: Homo sapiens (man)
C;Date: 17-Jan-1992 #sequence_revision 17-Jan-1992 #text_change 21-Jul-2000
C;Accession: A40036; S24543; A38142
R;Itoh, N.; Yonehara, S.; Ishii, A.; Yonehara, M.; Mizushima, S.I.; Sameshima, M.; Ha
Cell 66, 233-243, 1991
A;Title: The polypeptide encoded by the cDNA for human cell surface antigen Fas can m
A;Reference number: A40036; MUID:91309137
A;Accession: A40036
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-335 <ITO>
A;Cross-references: GB:M67454; NID:g182409; PIDN:AAA63174.1; PID:g182410
R;Krammer, P.H.
submitted to the EMBL Data Library, February 1992
A;Reference number: S24543
A;Accession: S24543
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-335 <KRA>
A;Cross-references: EMBL:X63717; NID:g28741; PID:g28742
R;Oehm, A.; Behrmann, I.; Falk, W.; Pawlita, M.; Maier, G.; Klas, C.; Li-Weber, M.; R
J. Biol. Chem. 267, 10709-10715, 1992
A;Title: Purification and molecular cloning of the APO-1 cell surface antigen, a memb
A;Reference number: A38142; MUID:92268122
A;Accession: A38142
A;Status: preliminary; not compared with conceptual translation
A;Molecule type: nucleic acid
A;Residues: 1-134,'Q',136-335 <OEH>
A;Experimental source: SKW6.4 cells
A;Note: sequence extracted from NCBI backbone (NCBIP:103810)

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A;Note: in NCBI backbone the source is designated as mouse
C;Genetics:
A;Gene: GDB:APT1
A;Cross-references: GDB:132671; OMIM:134637
A;Map position: 10q24.1-10q24.1
C;Superfamily: NGF receptor repeat homology
C;Keywords: apoptosis; surface antigen; transmembrane protein
F;1-16/Domain: signal sequence #status predicted <SIG>
F;85-128/Domain: NGF receptor repeat homology <NG4>
F;174-190/Domain: transmembrane #status predicted <TMM>


  Query Match             8.5%;  Score 197;  DB 2;  Length 335;
  Best Local Similarity   26.6%;  Pred. No. 9.6e-07;
  Matches   77;  Conservative   34;  Mismatches   93;  Indels   86;  Gaps   14;

Qy      38 LVLVVAAVLLL----VSAESALITQQDLAPQQRAAPQQKRSSPSEGL----------CPP 83
           | ||: :|  |    |:|:   |  : |  : |       :   |||          |||
Db       8 LPLVLTSVARLSSKSVNAQVTDINSKGL--ELRKTVTTVETQNLEGLHHDGQFCHKPCPP 65

Qy      84 GHH------ISEDGRDCISCKYGQDYSTHWNDLLFCLRCTRCDSG---EVELSPCTTTRN 134
           |        :: |  ||: |: |::|:   :    | ||  || |   |||:: || |:|
Db      66 GERKARDCTVNGDEPDCVPCQEGKEYTDKAHFSSKCRRCRLCDEGHGLEVEIN-CTRTQN 124

Qy     135 TVCQCEEGTFREEDSPEMCRKCRTGCPRGMVKVGDCTPWSDIECVHKESGT--------- 185
           | |:|:   |      | |  | | |  |::|  :||  |: :|  || |:
Db     125 TKCRCKPNFFCNSTVCEHCDPC-TKCEHGIIK--ECTLTSNTKC--KEEGSRSNLGWLCL 179

Qy     186 --------------------KHSGEAPAVEETVTSSPGTPA---SPCSLSGIIIGVTVA 221
                              || |     |: | :| | |   |   ||  |   |:|
Db     180 LLLPIPLIVWVKRKEVQKTCRKHRKENQGSHESPTLNPETVAINLSDVDLSKYI--TTIA 237

Qy     222 AVVLIVAVFVCKSLLWKKVLPYLKGICSGGGGDPERVDRSSQRPGAEDNV 271
           |:          | :||    | :  ::|       |||
Db     238 GVM--------------TLSQVKGFVRKNGVNEAKIDEIK-----NDNV 267


RESULT   7
A46484
apoptosis-mediating membrane-associated polypeptide Fas - mouse
C;Species: Mus musculus (house mouse)
C;Date: 18-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 05-Nov-1999
C;Accession: A46484; A47254
R;Watanabe-Fukunaga, R.; Brannan, C.I.; Itoh, N.; Yonehara, S.; Copeland, N.G.; Jenkins,
J. Immunol. 148, 1274-1279, 1992
A;Title: The cDNA structure, expression, and chromosomal assignment of the mouse Fas ant
A;Reference number: A46484; MUID:92148151
A;Accession: A46484
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-327 <WAT>
A;Cross-references: GB:M83649; NID:g193225; PIDN:AAA37593.1; PID:g193226
A;Experimental source: BAM3 macrophage cell line
A;Note: sequence extracted from NCBI backbone (NCBIN:81544, NCBIP:81545)
R;Adachi, M.; Watanabe-Fukunaga, R.; Nagata, S.
Proc. Natl. Acad. Sci. U.S.A. 90, 1756-1760, 1993
A;Title: Aberrant transcription caused by the insertion of an early transposable element
A;Reference number: A47254; MUID:93189576
A;Accession: A47254
A;Status: preliminary
A;Molecule type: nucleic acid
A;Residues: 1-96 <ADA>
A;Cross-references: GB:S56490; NID:g298505; PIDN:AAB25700.1; PID:g298506
A;Experimental source: MRL lpr/lpr
A;Note: sequence extracted from NCBI backbone (NCBIN:126850, NCBIN:126853, NCBIN:126863,
C;Superfamily: NGF receptor repeat homology
C;Keywords: transmembrane protein
F;44-79/Domain: NGF receptor repeat homology <NGF>
F;81-124/Domain: NGF receptor repeat homology <NG4>


  Query Match             7.7%;  Score 179.5;  DB 2;  Length 327;
  Best Local Similarity   24.4%;  Pred. No. 1.6e-05;
  Matches   87;  Conservative   49;  Mismatches  125;  Indels   95;  Gaps   20;

Qy      38 LVLVVAAVLLLVSAESALITQQDLAPQQRAAPQQKRSSPSEGL----------CPPGHHI 87
           | ||:|   | |  :      : |  ::|     |   ||||          | ||
Db       8 LPLVLAGSQLRVHTQGTNSISESLKLRRRVHETDKNC--SEGLYQGGPFCCQPCQPGKKK 65

Qy      88 SEDGR------DCISCKYGQDY--STHWNDLLFCLRCTRCDSG---EVELSPCTTTRNTV 136
           || :       |  |  |::|    |: |   | ||| ||     ||| : || |:||
Db      66 VEDCKMNGGTPTCAPCTEGKEYMDKNHYADK--CRRCTLCDEEHGLEVETN-CTLTQNTK 122

Qy     137 CQCEEGTFREEDSPEMCRKCRTGCPRGMVKVGDCTPWSDIECVHKESGTKHSGEAPAVEE 196
           |:|:   : :    | | :|   |  | ::   ||  |:  |  |:|
Db     123 CKCKPDFYCDSPGCEHCVRC-ASCEHGTLE--PCTATSNTNC-RKQS------------- 165

Qy     197 TVTSSPGTPASPCSLSGIIIGVTVAAVVLIVAVFVC----KSLLWKKVLPYLKGICSGGG 252
                 | :   |  |:          |:||  ||:      |   ||:
Db     166 --------PRNRLWLLTIL-------VLLIPLVFIYRKYRKRKCWKR-----------RQ 199

Qy     253 GDPERVDRSSQRPGAEDNVLNEIVSILQPTQVPE--QEMEVQEPAEPTGVNMLSPGESEH 310
           |||   |:| |      |  |  :|      :|   ::| :||  :    | :  |: :
Db     200 DDPE--SRTSSRETIPMNASNLSLS----KYIPRIAEDMTIQEAKKFARENNIKEGKIDE 253

Qy     311 LLEPAEAERSQRR-RLLV----PANEGDPTETL---------RQCFDDFADLVPFD 352
           ::  :  : :::: :||:       : |  : |          |:  | | |:|  |
Db     254 IMHDSIQDTAEQKVQLLLCWYQSHGKSDAYQDLIKGLKKAECRRTLDKFQDMVQKD 309


RESULT   8
GQRTT1
tumor necrosis factor receptor 1 precursor - rat
N;Contains: tumor necrosis factor binding protein 1 (TNF blocking factor)
C;Species: Rattus norvegicus (Norway rat)
C;Date: 30-Jun-1992 #sequence_revision 07-Oct-1994 #text_change 22-Jun-1999
C;Accession: B36555
R;Himmler, A.; Maurer-Fogy, I.; Kroenke, M.; Scheurich, P.; Pfizenmaier, K.; Lantz, M
DNA Cell Biol. 9, 705-715, 1990
A;Title: Molecular cloning and expression of human and rat tumor necrosis factor rece
A;Reference number: A36555; MUID:91090841
A;Accession: B36555
A;Molecule type: mRNA
A;Residues: 1-461 <HIM>
A;Cross-references: GB:M63122; NID:g207361; PIDN:AAA42256.1; PID:g207362
C;Comment: This protein is one of two known receptors for both TNF-alpha (cachectin)
C;Superfamily: tumor necrosis factor receptor type 1; NGF receptor repeat homology
C;Keywords: duplication; glycoprotein; receptor; transmembrane protein
F;1-29/Domain: signal sequence #status predicted <SIG>
F;30-461/Product: tumor necrosis factor receptor type 1 #status predicted <MAT>
F;30-211/Domain: extracellular #status predicted <EXT>
F;30-201/Product: tumor necrosis factor binding protein #status predicted <TBP>
F;44-82/Domain: NGF receptor repeat homology <NG1>
F;84-126/Domain: NGF receptor repeat homology <NG2>
F;127-167/Domain: NGF receptor repeat homology <NG3>
F;168-204/Domain: NGF receptor repeat homology <NG4>
F;212-234/Domain: transmembrane #status predicted <MEM>
F;235-461/Domain: intracellular #status predicted <INT>
F;54,151,201/Binding site: carbohydrate (Asn) (covalent) #status predicted


  Query Match             7.5%;  Score 173.5;  DB 1;  Length 461;
  Best Local Similarity   21.4%;  Pred. No. 6.4e-05;
  Matches  107;  Conservative   53;  Mismatches  168;  Indels  171;  Gaps   25;

Qy      32 PRVPKTLVLVVAAVLLL---VSAESALI-TQQDLAPQQRAAPQQKRSSPSEG-----LCP 82
           | ||  |: :|   ||:    |  : |: :  |   :    || | : |         |
Db       4 PIVPGLLLSLVLLALLMGIHPSGVTGLVPSLGDREKRDNLCPQGKYAHPKNNSICCTKCH 63

Qy      83 PGHHISED----GRD--CISCKYGQDYSTHWNDLLFCLRCTRC--DSGEVELSPCTTTRN 134
           | ::  |     |::  |  |  |  ::   | :  || |  |  :  :||:|||    :
Db      64 KGTYLVSDCPSPGQETVCEVCDKG-TFTASQNHVRQCLSCKTCRKEMFQVEISPCKADMD 122

Qy     135 TVCQCEEGTFRE--EDSPEMCRKCRTGCPRGMVKV-------------------GDCTP 172
           ||| |::  |:    ::   |  | : |  | | :                      :|||
Db     123 TVCGCKKNQFQRYLSETHFQCVDC-SPCFNGTVTIPCKEKQNTVCNCHAGFFLSGNECTP 181
```

```
Qy     173 WSDIECVHKESGTKHSGEA------PAVEETVTSSPGTPASPCSLSGIIIGVTVAAVVLI 226
Db     182 -----CSH----CKKNQECMKLCLPPVANVTNPQDSGT---------AVLLPLVIFLGLC 223

Qy     227 VAVFVCKSLL-----WKKVLPYLKGICSGGGGDPERVDRSSQRPGAEDNVLNEIVSILQP 281
Db     224 LLFFICISLLCRYPQW-------------------------RP--------RVYSIICR 249

Qy     282 TQVPEQEME----VQEPAEPTGVNMLSPGE-----------------SEHLLEPAEAER 319
Db     250 DSAPVKEVEGEGIVTKPLTPASIPAFSPNPGFNPTLGFSTTPRFSHPVSSTPISPVFGPS 309

Qy     320 SQRR-----RLLVPANEGDP-----------TETLRQCFDDFA----------------- 346
Db     310 NWHNFVPPVREVVPTQGADPLLYGSLNPVPIPAPVRKWEDVVAAQPQRLDTADPAMLYAV 369

Qy     347 -DLVPFDSWEPLMRKLGLMDNEI-KVAKAEAAGHRDTLYTMLIKWVNKTGR-----DASV 399
Db     370 VDGVPPTRWKEFMRLLGLSEHEIERLELQNGRCLREAHYSMLEAWRRRTPRHEATLDVVG 429

Qy     400 HTLLD-----ALETLGERL 413
Db     430 RVLCDMNLRGCLENIRETL 448
```

RESULT 9
A35356
tumor necrosis factor receptor 2 precursor [validated] - human
N;Alternate names: 75K tumor necrosis factor receptor; TNF receptor type 2
C;Species: Homo sapiens (man)
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 08-Dec-2000
C;Accession: A35356; A36475; A48416; A36007; A23666; B35010; I38094
R;Smith, C.A.; Davis, T.; Anderson, D.; Solam, L.; Beckmann, M.P.; Jerzy, R.; Dower, S.K
Science 248, 1019-1023, 1990
A;Title: A receptor for tumor necrosis factor defines an unusual family of cellular and
A;Reference number: A35356; MUID:90260639
A;Accession: A35356
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-461 <SMI>
A;Cross-references: GB:M32315; NID:g189185; PIDN:AAA59929.1; PID:g189186
R;Kohno, T.; Brewer, M.T.; Baker, S.L.; Schwartz, P.E.; King, M.W.; Hale, K.K.; Squires,
Proc. Natl. Acad. Sci. U.S.A. 87, 8331-8335, 1990
A;Title: A second tumor necrosis factor receptor gene product can shed a naturally occur
A;Reference number: A36475; MUID:91045991
A;Accession: A36475
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-195,'R',197-461 <KOH>
A;Cross-references: GB:M55994; GB:M38549; NID:g339757; PIDN:AAA36755.1; PID:g339758
R;Dembic, Z.; Loetscher, H.; Gubler, U.; Pan, Y.C.; Lahm, H.W.; Gentz, R.; Brockhaus, M.
Cytokine 2, 231-237, 1990
A;Title: Two human TNF receptors have similar extracellular, but distinct intracellular,
A;Reference number: A48416; MUID:91370690
A;Accession: A48416
A;Status: preliminary
A;Molecule type: mRNA; protein
A;Residues: 23-461 <DEM>
A;Cross-references: GB:S63368; NID:g235648; PIDN:AAB19824.1; PID:g235649
A;Note: sequence extracted from NCBI backbone (NCBIN:63368, NCBIP:63371)
R;Heller, R.A.; Song, K.; Onasch, M.A.; Fischer, W.H.; Chang, D.; Ringold, G.M.
Proc. Natl. Acad. Sci. U.S.A. 87, 6151-6155, 1990
A;Title: Complementary DNA cloning of a receptor for tumor necrosis factor and demonstra
A;Reference number: A36007; MUID:90349572
A;Accession: A36007
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 116-140,'P',142-195,'R',197-362,'T',364-461 <HEL>
A;Cross-references: GB:M35857; NID:g339751; PIDN:AAA63262.1; PID:g339752
R;Loetscher, H.; Schlaeger, E.J.; Lahm, H.W.; Pan, Y.C.E.; Lesslauer, W.; Brockhaus, M.
J. Biol. Chem. 265, 20131-20138, 1990
A;Title: Purification and partial amino acid sequence analysis of two distinct tumor ned
A;Reference number: A23666; MUID:91056048
A;Accession: A23666
A;Status: preliminary
A;Molecule type: protein
A;Residues: 23-40;65-69;136-141;300-306 <LOE>
R;Engelmann, H.; Novick, D.; Wallach, D.
J. Biol. Chem. 265, 1531-1536, 1990
A;Title: Two tumor necrosis factor-binding proteins purified from human urine. Eviden
A;Reference number: A35010; MUID:90110215
A;Accession: B35010
A;Status: preliminary
A;Molecule type: protein
A;Residues: 27-31 <ENG>
R;Kuhnert, P.; Kemper, O.; Wallach, D.
Gene 150, 381-386, 1994
A;Title: Cloning, sequencing and partial functional characterization of the 5' region
A;Reference number: I38094; MUID:95121934
A;Accession: I38094
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-37 <RES>
A;Cross-references: EMBL:X80021; NID:g666044; PIDN:CAA56324.1; PID:g825701
C;Genetics:
A;Gene: GDB:TNFR2
A;Cross-references: GDB:125914; OMIM:191191
A;Map position: 1p36.2-1p36.2
A;Introns: 26/3
A;Note: the list of introns is incomplete
C;Superfamily: tumor necrosis factor receptor type 2; NGF receptor repeat homology
C;Keywords: duplication; glycoprotein; receptor; transmembrane protein
F;1-22/Domain: signal sequence #status predicted <SIG>
F;23-416/Product: tumor necrosis factor receptor 2 #status experimental <MAT>
F;40-76/Domain: NGF receptor repeat homology <NG1>
F;78-119/Domain: NGF receptor repeat homology <NG2>
F;120-162/Domain: NGF receptor repeat homology <NG3>
F;164-201/Domain: NGF receptor repeat homology <NG4>
F;262-279/Domain: transmembrane #status predicted <TMN>
F;280-461/Domain: intracellular #status predicted <INT>
F;171,193/Binding site: carbohydrate (Asn) (covalent) #status predicted

Query Match 7.4%; Score 172; DB 1; Length 461;
Best Local Similarity 22.4%; Pred. No. 8.1e-05;
Matches 88; Conservative 40; Mismatches 134; Indels 130; Gaps 17;

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Qy      81 CPPGHH-----ISEDGRDCISCKYGQDYSTHWNDLLFCLRC-TRCDSGEVELSPCTTTRN 134
Db      57 CSPGQHAKVFCTKTSDTVCDSCE-DSTYTQLWNWVPECLSCGSRCSSDQVETQACTREQN 115

Qy     135 TVCQCEEGTFREEDSPEMC------RKCRTGCPRGMVKVGDCTPWSDIECVHKESGT--- 185
Db     116 RICTCRPGWYCALSKQEGCRLCAPLRKCRPGF--GVARPG--TETSDVVCKPCAPGTFSN 171

Qy     186 ------------------------------------------------------KHSGEAP 192
Db     172 TTSSTDICRPHQICNVVAIPGNASMDAVCTSTSPTRSMAPGAVHLPQPVSTRSQHTQPTP 231

Qy     193 AVEETVTSS---PGTPASPCSLS--------GIIIGVTVAAVVLIVAVFVC--KSLLWKK 239
Db     232 EPSTAPSTSFLLPMGPSPPAEGSTGDFALPVGLIVGVT-ALGLLIIGVVNCVIMTQVKKK 290

Qy     240 VL--------PYLKGICSGGGGDPERVDRSSQRPGAEDNVLNEIVSILQ---PTQVPEQE 288
Db     291 PLCLQREAKVPHLPADKARGTQGPEQQHLLITAPSSSSSSLESSASALDRRAPTRNQPQA 350

Qy     289 MEVQ-----EPAEPTGVNMLSPG---------------ESEHLLEPAEAERSQRRRLLV 327
Db     351 PGVEASGAGEARASTGSSDSSPGGHGTQVNVTCIVNVCSSSDH-----SSQCSSQASSTM 405

Qy     328 PANEGDPTETLRQCFDDFADLVPFDSWEPLMR 359
Db     406 GDTDSSPSESPKD------EQVPFSKEECAFR 431
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```
RESULT  10
I48854
gene murine tumour necrosis factor receptor 2 protein - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 02-Jul-1996 #sequence_revision 02-Jul-1996 #text_change 23-Jul-1999
C;Accession: I48854
R;Powell, E.E.; Wicker, L.S.; Peterson, L.B.; Todd, J.A.
Mamm. Genome 5, 726-727, 1994
A;Title: Allelic variation of the type 2 tumor necrosis factor receptor gene.
A;Reference number: I48854; MUID:95178848
A;Accession: I48854
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-459 <RES>
A;Cross-references: EMBL:X76401; NID:g433830; PIDN:CAA53981.1; PID:g433831
C;Superfamily: tumor necrosis factor receptor type 2; NGF receptor repeat homology
F;151-188/Domain: NGF receptor repeat homology <NGF>


  Query Match             7.2%;  Score 167.5;  DB 2;  Length 459;
  Best Local Similarity   22.0%;  Pred. No. 0.00017;
  Matches   71;  Conservative   30;  Mismatches  127;  Indels   95;  Gaps   13;

Qy      81 CPPGHHISE-----DGRDCISCKYGQDYSTHWNDLLFCLRC-TRCDSGEVELSPCTTTRN 134
           |||| ::          |  |:    |:  ||    || | : | : :||   ||  :|
Db      43 CPPGQYVKHFCNKTSDTVCADCEASM-YTQVWNQFRTCLSCSSSCSTDQVETRACTKQQN 101

Qy     135 TVCQCEEGTF----REEDSPEMC---RKCRTG-------CPRGMVKVGDCTPW------- 173
            || || | :        |   |    ||  |         | | |    | |
Db     102 RVCACEAGRYCALKTHSGSCRQCMRLSKCGPGFGVASSRAPNGNVLCKACAPGTFSDTTS 161

Qy     174 --------------------SDIECVHKESGTKH--------SGEAPAVEETVTSSPGT 204
                               :|  |   || |          |   |   : :   ||
Db     162 STDVCRPHRICSILAIPGNASTDAVCA-PESPTLSAIPRTLYVSQPEPTRSQPLDQEPGP 220

Qy     205 PASPCSLS--------------------GIIIGVTVAAVVLIVAVFVCKSLLWKKVLPYL 244
             :|  |:                    |:|:||| :  :|:: :  |  |: :|
Db     221 SQTPSILTSLGSTPIIEQSTKGGISLPIGLIVGVT-SLGLLMLGLVNCFILVQRK----- 274

Qy     245 KGICSGGGGDPERVDRSSQRPGAEDNVLNEIVSILQP---TQVPEQEMEVQEPAEPTGVN 301
                   |  : | :: |   |    : | : |    | |      | :   |
Db     275 --------KKPSCLQRDAKVPHVPDEKSQDAVGLEQQHLLTTAPSSSSSSLESSASAGDR 326

Qy     302 MLSPGESEHLLEPAEAERSQRRR 324
           ||        |||: ||  |
Db     327 RAPPGGHPQARVMAEAQGSQEAR 349


RESULT  11
GQHUN
nerve growth factor receptor precursor, low affinity [validated] - human
N;Alternate names: NGF receptor
C;Species: Homo sapiens (man)
C;Date: 31-Mar-1988 #sequence_revision 31-Mar-1988 #text_change 08-Dec-2000
C;Accession: A25218; A60204; S21689; I57638
R;Johnson, D.; Lanahan, A.; Buck, C.R.; Sehgal, A.; Morgan, C.; Mercer, E.; Bothwell, M.
Cell 47, 545-554, 1986
A;Title: Expression and structure of the human NGF receptor.
A;Reference number: A25218; MUID:87051725
A;Accession: A25218
A;Molecule type: mRNA
A;Residues: 1-427 <JOH>
A;Cross-references: GB:M14764; NID:g189204; PIDN:AAB59544.1; PID:g189205
R;Marano, N.; Dietzschold, B.; Earley Jr., J.J.; Schatteman, G.; Thompson, S.; Grob, P.;
J. Neurochem. 48, 225-232, 1987
A;Title: Purification and amino terminal sequencing of human melanoma nerve growth factc
A;Reference number: A60204; MUID:87085574
A;Accession: A60204
A;Molecule type: protein
A;Residues: 29-31,'T',33-42,'TT',45-46,'TX',50-51,'XX',54-56 <MAR>
A;Experimental source: melanoma cell line A875
A;Note: this sequence has been corrected by a note added in proof to follow the nucleoti
R;Vissavajjhala, P.; Leszyk, J.D.; Lin-Goerke, J.; Ross, A.H.
Arch. Biochem. Biophys. 294, 244-252, 1992
A;Title: Structural domains of the extracellular domain of human nerve growth factor
A;Reference number: S21689; MUID:92198017
A;Accession: S21689
A;Status: preliminary
A;Molecule type: protein
A;Residues: 183-208 <VIS>
R;Sehgal, A.; Patil, N.; Chao, M.
Mol. Cell. Biol. 8, 3160-3167, 1988
A;Title: A constitutive promoter directs expression of the nerve growth factor recept
A;Reference number: I57638; MUID:89096903
A;Accession: I57638
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-22 <RES>
A;Cross-references: GB:M21621; NID:g189206; PIDN:AAA36363.1; PID:g189207
C;Comment: This receptor is found on sensory and sympathetic neurons, on neuroblastom
C;Comment: The cysteine-rich region of the extracellular domain may form part or all
C;Comment: This protein is thought to form a high-affinity receptor when it associate
C;Comment: This receptor undergoes both N- and O-linked glycosylation.
C;Genetics:
A;Gene: GDB:NGFR
A;Cross-references: GDB:120234; OMIM:162010
A;Map position: 17q21-17q22
C;Superfamily: nerve growth factor receptor; NGF receptor repeat homology
C;Keywords: duplication; glycoprotein; heterodimer; monomer; phosphoprotein; receptor
F;1-28/Domain: signal sequence #status predicted <SIG>
F;29-427/Product: nerve growth factor receptor #status experimental <MAT>
F;29-250/Domain: extracellular #status predicted <EXT>
F;32-65/Domain: NGF receptor repeat homology <NG1>
F;67-108/Domain: NGF receptor repeat homology <NG2>
F;109-147/Domain: NGF receptor repeat homology <NG3>
F;149-189/Domain: NGF receptor repeat homology <NG4>
F;197-248/Region: serine/threonine-rich
F;251-272/Domain: transmembrane #status predicted <TRM>
F;273-427/Domain: intracellular #status predicted <INT>
F;60/Binding site: carbohydrate (Asn) (covalent) #status predicted


  Query Match             7.1%;  Score 166;  DB 1;  Length 427;
  Best Local Similarity   20.0%;  Pred. No. 0.0002;
  Matches   98;  Conservative   54;  Mismatches  167;  Indels  170;  Gaps   20;

Qy      28 ARPGPRVPKTLVLVVAAVLLLVSAESALITQQDLAPQQRAAPQQKRSSPSEGLCPPGHHI 87
           |  |||:   |:| |:    |  |: |                              || | :
Db       9 AMDGPRLLLLLLLGVS----LGGAKEA-------------------------CPTGLY- 37

Qy      88 SEDGRDCISCKYGQDY---------------STHWNDLLF----CLRCTRCDSGEVELS 127
           :  |  | :|  |:                  |  ::|::      |  || |    :   :
Db      38 THSGECCKACNLGEGVAQPCGANQTVCEPCLDSVTFSDVVSATEPCKPCTECVGLQSMSA 97

Qy     128 PCTTTRNTVCQCEEGTFREEDSP--EMCRKCRTG----------------CPRGMV--- 165
           ||    : ||:|  | :::| :   | || |  |                 || |
Db      98 PCVEADDAVCRCAYGYYQDETTGRCEACRVCEAGSGLVFSCQDKQNTVCEECPDGTYSDE 157

Qy     166 -----------------KVGDCTPWSDIECVH---------------KESGTKHSGEA 191
                            :: :|| |:| ||                 : :    ||
Db     158 ANHVDPCLPCTVCEDTERQLRECTRWADAECEEIPGRWITRSTPPEGSDSTAPSTQEPEA 217

Qy     192 PAVEETVTSSPG--------------TPASPCSLSGIIIGVTVAAVVLIVAVFVCKSLLW 237
           |  :: : |:                |  :  :|  :   :  | || :||    |   |
Db     218 PPEQDLIASTVAGVVTTVMGSSQPVVTRGTTDNLIPVYCSILAAVVVGLVAYIAFKR--W 275

Qy     238 KKVLPYLKGICSGGGGDPERVDRSSQRPGAEDNVLNEIVSILQPTQVPEQEMEVQEPAEP 297
                |    |     |       :| |  |   |:    |      |  | :  |:|
Db     276 NS----CKQNKQGANSRP-----VNQTPPPEGEKLHSDSGI----SVDSQSLHDQQPHTQ 322

Qy     298 T--GVNMLSPGESEHLLEPAEAERSQRRRLLVPANEGDPTETLRQCFDDFADLVPFDSWE 355
           |  |  :   |      | ||: |  ::   |:  : |                    |:|
Db     323 TASGQALKGDGGLYSSLPPAKREEVEK---LLNGSAG-----------------DTWR 360

Qy     356 PLMRKLGLMDNEIKVAKAEAAGHRDTLYTMLIKWVNKTGRDASVHTLLDALETLGERLAK 415
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Db    361 HLAGELGYQPEHIDSFTHEACPVR----ALLASWA--TQDSATLDALLAAL----RRIQR 410

Qy    416 QKIEDHLLS 424
Db    411 ADLVESLCS 419
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RESULT 12
B38634
tumor necrosis factor receptor type 2 precursor - mouse
C;Species: Mus musculus (house mouse)
C;Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 23-Jul-1999
C;Accession: B38634; A40254; S54816
R;Lewis, M.; Tartaglia, L.A.; Lee, A.; Bennett, G.L.; Rice, G.C.; Wong, G.H.W.; Chen, E.
Proc. Natl. Acad. Sci. U.S.A. 88, 2830-2834, 1991
A;Title: Cloning and expression of cDNAs for two distinct murine tumor necrosis factor r
A;Reference number: A38634; MUID:91187885
A;Accession: B38634
A;Molecule type: mRNA
A;Residues: 1-474 <LEW>
A;Cross-references: GB:M60469; NID:g199827; PIDN:AAA39752.1; PID:g199828
R;Goodwin, R.G.; Anderson, D.; Jerzy, R.; Davis, T.; Brannan, C.I.; Copeland, N.G.; Jenk
Mol. Cell. Biol. 11, 3020-3026, 1991
A;Title: Molecular cloning and expression of the type 1 and type 2 murine receptors for
A;Reference number: A40254; MUID:91246168
A;Accession: A40254
A;Molecule type: mRNA
A;Residues: 1-474 <GOO>
A;Cross-references: GB:M60469; NID:g199827; PIDN:AAA39752.1; PID:g199828
R;Kissonerghis, M.; Fellowes, R.; Feldmann, M.; Chernajovsky, Y.
submitted to the EMBL Data Library, May 1995
A;Description: Characterization of the promoter region of the murine p75-TNF receptor.
A;Reference number: S54816
A;Accession: S54816
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-22 <KIS>
A;Cross-references: EMBL:X87128; NID:g809043; PIDN:CAA60618.1; PID:g809044
C;Superfamily: tumor necrosis factor receptor type 2; NGF receptor repeat homology
C;Keywords: cytokine receptor; transmembrane protein
F;1-22/Domain: signal sequence #status predicted <SIG>
F;23-474/Product: tumor necrosis factor receptor type 2 #status predicted <MAT>
F;40-77/Domain: NGF receptor repeat homology <NG1>
F;79-120/Domain: NGF receptor repeat homology <NG2>
F;166-203/Domain: NGF receptor repeat homology <NG4>

Query Match 7.0%; Score 163.5; DB 2; Length 474;
Best Local Similarity 21.7%; Pred. No. 0.00034;
Matches 70; Conservative 31; Mismatches 127; Indels 95; Gaps 13;

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Qy     81 CPPGHHISE-----DGRDCISCKYGQDYSTHWNDLLFCLRC-TRCDSGEVELSPCTTTRN 134
Db     58 CPPGQYVKHFCNKTSDTVCADCEASM-YTQVWNQFRTCLSCSSSCTTDQVEIRACTKQQN 116

Qy    135 TVCQCEEGTF----REEDSPEMC---RKCRTG-------CPRGMVKVGDCTPW------- 173
Db    117 RVCACEAGRYCALKTHSGSCRQCMRLSKCGPGFGVASSRAPNGNVLCKACAPGTFSDTTS 176

Qy    174 --------------------SDIECVHKESGTKH--------SGEAPAVEETVTSSPGT 204
Db    177 STDVCRPHRICSILAIPGNASTDAVCA-PESPTLSAIPRTLYVSQPEPTRSQPLDQEPGP 235

Qy    205 PASPCSLS--------------------GIIIGVTVAAVVLIVAVFVCKSLLWKKVLPYL 244
Db    236 SQTPSILTSLGSTPIIEQSTKGGISLPIGLIVGVT-SLGLLMLGLVNCIILVQRK----- 289

Qy    245 KGICSGGGGDPERVDRSSQRPGAEDNVLNEIVSILQP---TQVPEQEMEVQEPAEPTGVN 301
Db    290 --------KKPSCLQRDAKVPHVPDEKSQDAVGLEQQHLLTTAPSSSSSSLESSASAGDR 341

Qy    302 MLSPGESEHLLEPAEAERSQRRR 324
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Db    342 RAPPGGHPQARVMAEAQGFQEAR 364
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RESULT 13
JN0006
nerve growth factor receptor, low affinity precursor - chicken
N;Alternate names: NGF receptor
C;Species: Gallus gallus (chicken)
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 10-Sep-1999
C;Accession: JN0006; A60504
R;Large, T.H.; Weskamp, G.; Helder, J.C.; Radeke, M.J.; Misko, T.P.; Shooter, E.M.; R
Neuron 2, 1123-1134, 1989
A;Title: Structure and developmental expression of the nerve growth factor receptor i
A;Reference number: JN0006; MUID:90166579
A;Accession: JN0006
A;Molecule type: mRNA
A;Residues: 1-416 <LAR>
A;Experimental source: embryonic chick brain
R;Heuer, J.G.; Fatemie-Nainie, S.; Wheeler, E.F.; Bothwell, M.
Dev. Biol. 137, 287-304, 1990
A;Title: Structure and developmental expression of the chicken NGF receptor.
A;Reference number: A60504; MUID:90152140
A;Accession: A60504
A;Status: preliminary; not compared with conceptual translation
A;Molecule type: mRNA
A;Residues: 21-35,'Y',37-172,'K',174-275,'S',277-395,'R',397-416 <HEU>
C;Comment: This receptor is found on sensory and sympathetic neurons, on neuroblastom
C;Comment: The cysteine-rich region of the extracellular domain may form part or all
C;Comment: This protein is thought to form a high-affinity receptor when it associate
C;Superfamily: nerve growth factor receptor; NGF receptor repeat homology
C;Keywords: duplication; glycoprotein; heterodimer; monomer; phosphoprotein; receptor
F;1-20/Domain: signal sequence #status predicted <SIG>
F;21-416/Product: nerve growth factor receptor #status predicted <MAT>
F;21-239/Domain: extracellular #status predicted <EXT>
F;24-57/Domain: NGF receptor repeat homology <NG1>
F;59-100/Domain: NGF receptor repeat homology <NG2>
F;101-139/Domain: NGF receptor repeat homology <NG3>
F;141-181/Domain: NGF receptor repeat homology <NG4>
F;189-237/Region: serine/threonine-rich
F;240-261/Domain: transmembrane #status predicted <MEM>
F;262-416/Domain: intracellular #status predicted <INT>
F;52/Binding site: carbohydrate (Asn) (covalent) #status predicted

Query Match 6.9%; Score 161.5; DB 1; Length 416;
Best Local Similarity 22.8%; Pred. No. 0.0004;
Matches 91; Conservative 46; Mismatches 160; Indels 103; Gaps 20;

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Qy     81 CPP-----GHH------ISEDGRDCISCKYGQDYSTHWNDLL--FCLRCTRCDSGEVELS 127
Db     75 CKPCTQCVGLHSMSAPCVESDDAVC-RCAYG-----YFQDELSGSCKECSICEVGFGLMF 128

Qy    128 PCTTTRNTVC-QCEEGTFREE-DSPEMCRKCRTGCPRGMVKVGDCTPWSDIEC--VHKES 183
Db    129 PCRDSQDTVCEECPEGTFSDEANFVDPCLPC-TICEENEVMVKECTATSDAECRDLHPR- 186

Qy    184 GTKHS--------------------GEAPAVEETVTSSPGTPASPCS------LSGIIIG 217
Db    187 WTTHTPSLAGSDSPEPITRDPFNTEGMATTLADIVTTVMGSSQPVVSRGTADNLIPVYCS 246

Qy    218 VTVAAVVLIVAVFVCKSLLWKKVLPYLKGICSGGGGDPERVDRSSQRPGAEDNVLNEIVS 277
Db    247 ILAAVVVGLVAYIAFKR--WNS----CKQNKQGANNRP-----VNQTPSPEGEKLHSDSG 295

Qy    278 ILQPTQVPEQEMEVQEPAEPTGVNMLSPGESEHLLEPAEAERSQRRRLLVPANEGDPTET 337
Db    296 I----SVDSQSLHDQQPPNQSTQGPAPKGDGSLYASLPPSKQEEVEKLLSSSAE------ 345

Qy    338 LRQCFDDFADLVPFDSWEPLMRKLGLMDNEIKVAKAEAAGHRDTLYTMLIKWVNKTGRDA 397
Db    346 -------------ETWRQLAGELGYKEDLIDCFTREESPAR----ALLADW------SA 381
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```
Qy    398 SVHTLLDALETLGERLAKQKIEDHLLSSGKFMYLEGNADS 437
Db    382 KETATLDALLVALRKIQRGDIAESL-------YSESTATS 414


RESULT  14
A26431
nerve growth factor receptor precursor, low affinity - rat
N;Alternate names: NGF receptor
C;Species: Rattus norvegicus (Norway rat)
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 10-Sep-1999
C;Accession: A26431; PH1229
R;Radeke, M.J.; Misko, T.P.; Hsu, C.; Herzenberg, L.A.; Shooter, E.M.
Nature 325, 593-597, 1987
A;Title: Gene transfer and molecular cloning of the rat nerve growth factor receptor.
A;Reference number: A26431; MUID:87115859
A;Accession: A26431
A;Molecule type: mRNA
A;Residues: 1-425 <RAD>
A;Cross-references: GB:X05137; NID:g56755; PIDN:CAA28783.1; PID:g56756
R;Metsis, M.; Timmusk, T.; Allikmets, R.; Saarma, M.; Persson, H.
Gene 121, 247-254, 1992
A;Title: Regulatory elements and transcriptional regulation by testosterone and retinoic
A;Reference number: PH1229; MUID:93077038
A;Accession: PH1229
A;Molecule type: DNA
A;Residues: 1-20 <MET>
A;Cross-references: GB:X61269
C;Comment: This receptor is found on sensory and sympathetic neurons, on neuroblastoma c
C;Comment: The cysteine-rich region of the extracellular domain may form part or all of
C;Comment: This protein is thought to form a high-affinity receptor when it associates w
C;Genetics:
A;Introns: 20/3
C;Superfamily: nerve growth factor receptor; NGF receptor repeat homology
C;Keywords: duplication; glycoprotein; heterodimer; monomer; phosphoprotein; receptor; t
F;1-29/Domain: signal sequence #status predicted <SIG>
F;30-425/Product: nerve growth factor receptor #status predicted <MAT>
F;30-251/Domain: extracellular #status predicted <EXT>
F;33-66/Domain: NGF receptor repeat homology <NG1>
F;68-109/Domain: NGF receptor repeat homology <NG2>
F;110-148/Domain: NGF receptor repeat homology <NG3>
F;150-190/Domain: NGF receptor repeat homology <NG4>
F;198-249/Region: serine/threonine-rich
F;252-273/Domain: transmembrane #status predicted <MEM>
F;274-425/Domain: intracellular #status predicted <INT>
F;61/Binding site: carbohydrate (Asn) (covalent) #status predicted


  Query Match             6.7%;  Score 155;  DB 1;  Length 425;
  Best Local Similarity   23.2%;  Pred. No. 0.0012;
  Matches   89;  Conservative   45;  Mismatches  161;  Indels   88;  Gaps   17;

Qy     71 QKRSSPSEGLCPPGHHISEDGRDCISCKYG--QDYSTHWNDLLFCLRCTRCDSGEVELSP 128
Db     94 QSMSAP----C-----VEADDAVC-RCAYGYYQDEETG-----HCEACSVCEVGSGLVFS 138

Qy    129 CTTTRNTVC-QCEEGTFREE-DSPEMCRKCRTGCPRGMVKVGDCTPWSDIECVHKESGTK 186
Db    139 CQDKQNTVCEECPEGTYSDEANHVDPCLPC-TVCEDTERQLRECTPWADAEC--EEIPGR 195

Qy    187 HSGEAPAVEETVTSSPGTPASPCSLSGIIIGVTVAAVVLIVAVFVCKSLLWKKVLPYLKG 246
Db    196 WIPRSTPPEGSDSTAPSTQEPEVPPEQDLVPSTVADMVTTVM----------------- 237

Qy    247 ICSGGGGDPERVDRSSQRPGAEDNVLNEIVSILQPTQVP----------EQEMEVQEPAE 296
Db    238 -----GSSQPVVTR-----GTTDNLIPVYCSILAAVVVGLVAYIAFKRWNSCKQNKQGAN 287

Qy    297 PTGVNMLSPGESEHL---------LEPAEAERSQRRRLLVPANEGDPT------ETLRQC 341
Db    288 SRPVNQTPPPEGEKLHSDSGISVDSQSLHDQQTHTQTASGQALKGDGNLYSSLPLTKR-- 345

Qy    342 FDDFADLVPFDSWEPLMRKLGLMDNEIKVAKAEAAGHRDTLYTMLIKWVNKTGRDASVHT 401
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```
Db    346 -EEVEKLLNGDTWRHLAGELGYQPEHIDSFTHEACPVR----ALLASW------GAQDSA 394

Qy    402 LLDALETLGERLAKQKIEDHLLS 424
Db    395 TLDALLAALRRIQRADIVESLCS 417


RESULT  15
JC2395
Fas antigen precursor - rat
C;Species: Rattus norvegicus (Norway rat)
C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 05-Nov-1999
C;Accession: JC2395; PC2246
R;Kimura, K.; Wakatsuki, T.; Yamamoto, M.
Biochem. Biophys. Res. Commun. 198, 666-674, 1994
A;Title: A variant mRNA species encoding a truncated from of Fas antigen in the rat l
A;Reference number: JC2395; MUID:94128114
A;Accession: JC2395
A;Molecule type: mRNA
A;Residues: 1-324 <KIM>
A;Cross-references: DDBJ:D26112; NID:g468486; PIDN:BAA05108.1; PID:d1005650; PID:g468
A;Experimental source: thymus
A;Accession: PC2246
A;Molecule type: mRNA
A;Residues: 1-62,'RFT' <KI2>
A;Cross-references: DDBJ:D26113; NID:g468488; PIDN:BAA05109.1; PID:d1005651; PID:g468
A;Experimental source: liver
C;Genetics:
A;Introns: 62/1
C;Superfamily: NGF receptor repeat homology
C;Keywords: transmembrane protein
F;1-21/Domain: signal sequence #status predicted <SIG>
F;22-324/Product: Fas antigen #status predicted <MAT>
F;44-79/Domain: NGF receptor repeat homology <NGF>
F;81-124/Domain: NGF receptor repeat homology <NG4>
F;171-188/Domain: transmembrane #status predicted <TMM>


  Query Match             6.4%;  Score 148.5;  DB 2;  Length 324;
  Best Local Similarity   25.5%;  Pred. No. 0.0025;
  Matches   67;  Conservative   33;  Mismatches   84;  Indels   79;  Gaps   14;

Qy     39 VLVVAAVLLLVSAESALITQQDLAPQQRAAPQQKRS------SPSEGL----------CP 82
Db      1 MLWIMAVLPLVLAGPELNVRMQGTDSIFEGLELKRSVRETDNNCSEGLYQVGPFCCQPCQ 60

Qy     83 PGHHISED------GRDCISCKYGQDYS--THWNDLLFCLRCTRCDSG---EVELSPCTT 131
Db     61 PGERKVKDCTTSGGAPTCHPCTEGEEYTDRKHYSDK--CRRCAFCDEGHGLEVETN-CTR 117

Qy    132 TRNTVCQCEEGTFREEDSPEMCRKCRTGCPRGMVKVGD-CTPWSDIECVHKESGTKHSGE 190
Db    118 TQNTKCRCKENFYCNASLCDHCYHC-TSC--GLEDILEPCTRTSNTKCKKQSSNYK---- 170

Qy    191 APAVEETVTSSPGTPASPCSLSGIIIGVTVAAVVLIVAVFVCKSLLWKKVLPYLKGICSG 250
Db    171 ----------------------LLWLLILPGLAILFVFIYKR--YRKRQP-------- 196

Qy    251 GGGDPERVDRSSQRPGAEDNVLN 273
Db    197 --GDPE-----SGIPSPESVPMN 212


Search completed: May 23, 2001, 14:19:46
Job time: 91 sec
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